

PR 07-JUN-1995; 95US-0479799.

XX (THER-) THERA PRO.

PI Gasanov SE, Rael ED, Vernon LP.

DR WPI: 1997-065280/06.

DR N-PSDB: 747764.

PT New target specific toxins, partic for cancer cells - comprising a molecule capable of specific binding to the surface of a cell linked to pyruvate thionin peptide.

PS Claim 1; Page 36; 52pp; English.

CC This sequence is a Pyruvate thionin (PT) protein. Target specific CC toxins can be constructed by linking this toxin to a molecule (esp. CC monoclonal antibody anti-CD5) capable of specifically binding the surface of a cell. The target specific toxin can be used to kill selected CC undesirable cells to which PT is generally cytotoxic, partic. cancer CC cells. The immunotoxins can also be used for the manipulation of cells CC used in tissue and organ grafts, blood transfusions and bone marrow CC transplants and to treat graft-versus-host disease. The immunotoxins CC display a high degree of specificity and cytotoxicity. PT is membrane- CC active, obviating the need for PT to be internalised in order to exert CC its cytotoxic effect. PT is a very stable, compact peptide which is CC resistant to most proteases and is not immunogenic. The PT cytotoxicity is CC lost after it is incorporated into the lipid bilayer of a host cell so CC that it will not produce second round cytotoxicity towards macrophages CC and other cells that come in contact with the membrane of cells contg. CC the PT immunotoxin.

Sequence 48 AA:

Query Match 100.0%; Score 52; DB 18; Length 48;
Best Local Similarity 20.0%; Pred. No. 1.6e+02;
Matches 4; Conservative 16; Mismatches 0; Indels 0; Gaps 0;

Y 1 CXXXXXXXXXXXXXXCXXC 20
I:::|||||::|::|
b 13 cyovcrpglstrcicakc 32

ESULT 4
R6122 96122 standard; Peptide: 50 AA.

R96122:

17-DEC-1996 (first entry)

Leech derived fahsin based protease inhibitor #2.

Protease inhibitor; isoform; elastase; chymotrypsin; trypsin; leech;
tissue; secretion; saliva; fahsin; antibiotic; diabetes mellitus;
blood clotting disorder; neutrophil function; emphysema;
rheumatoid arthritis; HIV infection; human immunodeficiency virus.

Limnatis nilotica.

MO9613585-A1.

09-MAY-1996.

27-OCT-1995; 95MO-EP04223.

14-MAR-1995; 95EP-0103637.

28-OCT-1994; 94EP-0117053.

(CLOD-) CLODICA SA.

Voerman G;

XX WPI: 1996-239498/24.

PT New protease inhibitors from the leech Limnatis nilotica - for
PT treating, e.g. blood clotting disorders, HIV infection, diabetes
PT mellitus etc.

PS Claim 3; Page 26; 41pp; English.

CC The protease inhibitor peptide isoforms given in R96121-23 are
CC elastase/chymotrypsin- and trypsin inhibitors which may be isolated
CC from leech tissue or leech secretions, e.g. saliva. These peptides
CC belong to the family of leech derived substances named fahsin's which
CC also have an antibiotic effect. The fahsin family of proteins comprise
CC 50/51 amino acids and occur in various isoforms. These peptides are
CC useful in the treatment of diabetes mellitus, blood clotting disorders,
CC disorders of neutrophil function, e.g. emphysema, rheumatoid arthritis,
CC HIV infection and other immunological and inflammatory diseases.

Sequence 50 AA:

Query Match 100.0%; Score 52; DB 17; Length 50;
Best Local Similarity 20.0%; Pred. No. 1.7e+02;
Matches 4; Conservative 16; Mismatches 0; Indels 0; Gaps 0;

Y 1 CXXXXXXXXXXXXXXCXXC 20
I:::|||||::|::|
Db 27 clycpkxgfevdegcelpc 46

RESULT 5

R96123 96123 standard; Peptide: 50 AA.

R96123:

17-DEC-1996 (first entry)

Leech derived fahsin based protease inhibitor #3.

Protease inhibitor; isoform; elastase; chymotrypsin; trypsin; leech;
tissue; secretion; saliva; fahsin; antibiotic; diabetes mellitus;
blood clotting disorder; neutrophil function; emphysema;
rheumatoid arthritis; HIV infection; human immunodeficiency virus.

Limnatis nilotica.

MO9613585-A1.

09-MAY-1996.

27-OCT-1995; 95MO-EP04223.

14-MAR-1995; 95EP-0103637.

28-OCT-1994; 94EP-0117053.

(CLOD-) CLODICA SA.

Voerman G;

WPI: 1996-239498/24.

PT New protease inhibitors from the leech Limnatis nilotica - for
PT treating, e.g. blood clotting disorders, HIV infection, diabetes
PT mellitus etc.

PS Claim 3; Page 26; 41pp; English.

CC The protease inhibitor peptide isoforms given in R96121-23 are
CC elastase/chymotrypsin- and trypsin inhibitors which may be isolated
CC from leech tissue or leech secretions, e.g. saliva. These peptides
CC belong to the family of leech derived substances named fahsin's which

[illegible]

FT /label= Unknown
 FT Misc-difference 21
 FT /label= Unknown
 FT /note= "Xaa may be 10 amino acids in length; some
 FT amino acids may be absent"
 FT Misc-difference 23
 FT /label= Unknown
 FT Misc-difference 24
 FT /label= Unknown
 FT Misc-difference 25
 FT /label= Unknown
 FT Misc-difference 27
 FT /label= Unknown
 FT /note= "Xaa may be 7 amino acids in length; some
 FT amino acids may be absent"
 FT Misc-difference 29
 FT /label= Unknown
 FT /note= "Xaa may be 27 amino acids in length; some
 FT amino acids may be absent"
 FT Misc-difference 31
 FT /label= Unknown
 FT /note= "Xaa may be 13 amino acids in length; some
 FT amino acids may be absent"
 FT WO200021555-A1.
 XX
 XX PD 20-APR-2000.
 XX
 XX PF 13-OCT-1999: 99WO-US23640.
 XX
 XX PR 15-OCT-1998: 98US-0104355.
 XX
 XX PA (HARD) HARVARD COLLEGE.
 XX
 XX PI McMahon AP, Parr BA, Vaino S;
 XX
 XX DR WPI: 2000-317845/27.
 XX
 XX PT Contraceptive composition for inhibiting oocyte development in a female
 XX primate comprises a Wnt polypeptide antagonist
 XX
 XX PS Claim 12: Page 44; 57pp: English.
 XX
 XX CC The patent discloses a method of female primate contraception comprising
 XX administering an antagonist of a Wnt polypeptide, inhibiting oocyte
 XX development. Wnt polypeptides are useful for promotive maturation of an
 XX immature oocyte. Wnt polypeptides are also useful for increasing the
 XX number of mature oocytes and to enhance oocyte viability. The present
 XX peptide is a consensus sequence of Wnt antagonist which inhibits the
 XX physiological activity of a Wnt polypeptide. Antagonistic polypeptides
 XX may contain a cysteine-rich domain.
 XX
 XX SQ Sequence 31 AA;

Query Match 100.0%; Score 53; DB 21; Length 31;
 Best Local Similarity 66.7%; Pred. No. 1.1e+02;
 Matches 14; Conservative 7; Mismatches 0; Indels 0; Gaps 0;

OY 1 CXXXXXXXCCCCXXXXC 21
 I:::|||||:|||||
 Db 6 CCCCCCXXXXXXCCXXXXC 26

RESULT 2
 R11372
 ID R11372 standard; Protein: 40 AA.
 XX
 AC R11372;
 XX
 08-MAY-1991 (first entry)

1 generated by genomic meg-csp program.

XX
 KW Megakaryocyte colony stimulating factor; platelet deficiency;
 KW bleeding disorder.
 XX
 OS Homo sapiens.
 XX
 XX W09102003-A.
 XX
 XX PD 21-FEB-1991.
 XX
 XX PF 07-AUG-1990: 90WO-US04421.
 XX
 XX PR 29-JUN-1990: 90US-0546114.
 XX 08-AUG-1989: 89US-0390901.
 XX PR 28-DEC-1989: 89US-0457196.
 XX
 XX (GENE-) GENETICS INST INC.
 XX
 XX PI Gesner TG, Clark SC, Turner K, Hewick RM;
 XX
 XX DR WPI: 1991-073490/10.
 XX
 XX DR N-PSDB: 010580.
 XX
 XX PS New mega:karyocyte colony stimulating factor protein - regulates
 XX PT human haematopoiesis by stimulating growth and development of
 XX PT mega:karyocyte(s) in treatment of e.g. plastic anaemia
 XX
 XX Claim 3: Page 85; 204pp: English.
 XX
 XX CC The clone was isolated from a human placenta lambda phage DNA
 XX library. The sequence can be inserted into expression vectors for
 XX the prodn. of recombinant meg-CSF. The protein is used to treat
 XX CC bleeding disorders and platelet deficiencies.
 XX See also R10870, R10871 and R10872.
 XX
 XX SQ Sequence 40 AA;

Query Match 100.0%; Score 53; DB 12; Length 40;
 Best Local Similarity 19.0%; Pred. No. 1.4e+02;
 Matches 4; Conservative 17; Mismatches 0; Indels 0; Gaps 0;

OY 1 CXXXXXXXCCCCXXXXC 21
 I:::|||||:|||||
 Db 4 Ckgrfcsfgrccdcagc 24

RESULT 3
 W02648
 ID W02648 standard; pepide: 49 AA.
 XX
 AC W02648;
 XX
 XX DT 23-OCT-1996 (first entry)
 XX
 XX DE Mutant disintegrin amino acid sequence.
 XX
 XX KW Wild type: RGD motif; ecstatin; disintegrin; binding activity.
 XX
 XX OS Synthetic.
 XX
 XX FH Key Location/Qualifiers
 XX FT Domain 24..26
 XX FT /note= "RGD domain"

XX JP08157456-A.
 XX
 XX PD 18-JUN-1996.
 XX
 XX PF 30-NOV-1994: 94JP-0296474.
 XX
 XX PR 30-NOV-1994: 94JP-0296474.
 XX

(TANP-)TANPAKU KOGAKU KENKYUSHO KK.
 WPI: 1996-339190/34.
 A mutant disintegrin - contg. Cys residues flanking the RGD functional site to cyclise it
 Disclosure: Fig 1: 6pp; Japanese.
 This is the amino acid sequence of a mutated sequence surrounding the RGD peptide motif in ecstasin, a member of the disintegrin family. The amino acids immediately flanking the RGD motif were mutated to Cys residues in order to circularise the RGD motif. The peptides were synthesised using a peptide synthesiser and their integrin binding activities determined and compared.
 Sequence 49 AA:
 Query Match 100.0%; Score 53; DB 17; Length 49;
 Best Local Similarity 19.0%; Pred. No. 1.6e+02;
 Matches 4: Conservative 17; Mismatches 0; Indels 0; Gaps 0
 1 CXXXCXXXXXXXXXXXXXC 21
 1:::|:::|:::|:::|:::|
 7 ccrnckfkgclckrgrdc 27
 RESULT 4
 Y57813
 ID Y57813 standard; protein: 57 AA.
 AC Y57813:
 XX
 XX
 DT 22-MAR-2000 (first entry)
 XX
 DE Crab metallothionein Class I amino acid sequence.
 XX
 XX Metallothionein: metal recovery; remediation: heavy metal;
 KM precious metal; phytochelatin; green algae; Chlamydomonas reinhardtii.
 XX
 OS Eubrachyura.
 OS
 PN WO9960838-A1.
 PD 02-DEC-1999.
 XX
 PF 28-MAY-1999: 99WO-US12007.
 XX
 PR 28-MAY-1998: 98US-0087374.
 XX
 PA (OHIS) UNIV OHIO STATE RES FOUND.
 XX
 PI Sayre RT, Traina SJ.
 DR WPI: 2000-086646/07.
 XX
 PT Novel method for metal recovery, remediation and separation
 XX
 PS Disclosure: Page 6: 86pp; English.
 XX
 CC The present invention describes a transgenic algal cell (1) of the
 CC genus Chlamydomonas comprising reproducible genetic material comprising
 CC a nucleotide sequence capable of expressing chicken type I
 CC Metallothionein. Also described is a method of removing metal from
 CC an aqueous medium containing at least one dissolved or suspended
 CC metal. The transgenic algae are used for the selective separation of
 CC metals, particularly the separation of precious and desirable metals
 CC such as gold and uranium, from other metals such as cadmium, zinc and
 CC copper. The method can be used to facilitate the selective recovery of
 CC precious and rare metals from mineral sources where aqueous media can
 CC be used, such as in natural surface water flows, ground water and where
 CC water may have been polluted. The method is suitable for well defined,

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CC soil and water remediation arts, mining fields, and industrial  
CC engineering. The present sequence represents a Class I metalloprotein  
CC given in the present invention.  
XX  
SQ Sequence 57 AA:  
  
Query Match 100.0%; Score 53; DB 21; Length 57;  
Best Local Similarly 19.0%, Pred. No. 1.8e+02;  
Matches 4: Conservative 17; Mismatches 0; Indels 0; Gaps 0;  
OY 1 CXXXCXXXXXXXXXXXXXCCXXC 21  
|::|::|::|::|::|::|::|  
Db 33 csgcckankgcgcrctskpc 53  
  
RESULT 5  
Y76185  
ID Y76185 standard; Protein; 57 AA.  
AC Y76185;  
XX  
XX 23-MAR-2000 (first entry)  
DT  
DE Human secreted protein encoded by gene 62.  
KW Human; secreted protein; cancer; tumour; developmental abnormality;  
KW foetal deficiency; blood disorder; immune system disorder; inflammation;  
KW autoimmune disease; allergy; Alzheimer's disease; cognitive disorder;  
KW schizophrenia; arthritis; asthma; psoriasis; sepsis; skin disorder;  
KW atherosclerosis; diabetes; cardiovascular disorder; kidney disorder;  
KW digestive disorder; endocrine disorder; infection; AIDS; leukaemia;  
KW therapy.  
XX  
XX Homo sapiens.  
OS  
XX W09958660-A1.  
PN  
PD 18-NOV-1999.  
XX  
PF 06-MAY-1999; 99MO-US09847.  
XX  
PR 12-MAY-1998; 98US-0085093.  
PR 12-MAY-1998; 98US-0085094.  
PR 12-MAY-1998; 98US-0085105.  
PR 12-MAY-1998; 98US-0085180.  
PR 18-MAY-1998; 98US-0085906.  
PR 18-MAY-1998; 98US-0085920.  
PR 18-MAY-1998; 98US-0085921.  
PR 18-MAY-1998; 98US-0085922.  
PR 18-MAY-1998; 98US-0085923.  
PR 18-MAY-1998; 98US-0085924.  
PR 18-MAY-1998; 98US-0085928.  
PR 18-MAY-1998; 98US-0085925.  
PPR 18-MAY-1998; 98US-0085927.  
XX  
PA (HUMA-) HUMAN GENOME SCI INC.  
XX  
PI Ruben SM, Florence K, Ni J, Rosen CA, Carter KC, Moore PA;  
PI Olsen HS, Shi Y, Young PE, Wei F, Brewer LA, Soppet DR;  
PI Lafleur DM, Endress CA, Ebner R;  
XX  
XX MPI: 2000-062296/05.  
DR N-PSDB; 265311.  
XX  
XX Claim 11; Page 398; 475pp; English.  
XX  
XX %6250 to %6250 represent 97 isolated human secreted protein genes.  
XX V01234 to V01234 represent the secreted proteins encoded by the 97 human
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